

**Atherosclerosis Risk in Communities (ARIC) Study
dbGaP Acknowledgment Statements
9-8-2011**

The acknowledgment of the ARIC dataset(s) should always include acknowledgment of the ARIC contract (component I below) **AND** the genotypic/genomic dataset(s) (component II below).

Components of the ARIC dbGaP Acknowledgment Statements:

I. Atherosclerosis Risk in Communities (ARIC) Contract Acknowledgment (Use the following statement for **ALL** analyses resulting from ARIC dbGaP data).

“The Atherosclerosis Risk in Communities Study is carried out as a collaborative study supported by National Heart, Lung, and Blood Institute contracts (HHSN268201100005C, HHSN268201100006C, HHSN268201100007C, HHSN268201100008C, HHSN268201100009C, HHSN268201100010C, HHSN268201100011C, and HHSN268201100012C). The authors thank the staff and participants of the ARIC study for their important contributions.”

II. Genotypic/Genomic Dataset Acknowledgment (Use one or more of the following statements relevant to the specific datasets used in your analysis)

A. ARIC Candidate gene Association Resource (CARE): “Funding for CARE genotyping was provided by NHLBI Contract N01-HC-65226.”

B. ARIC Gene Environment Association Studies (GENEVA): “Funding for GENEVA was provided by National Human Genome Research Institute grant U01HG004402 (E. Boerwinkle).”

C. CALiCo-ARIC Population Architecture of Genes and Environment(PAGE) Consortium

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D. NHLBI GO-ESP: Heart Cohorts Exome Sequencing Project (ARIC)

“This study is part of the NHLBI Grand Opportunity Exome Sequencing Project (GO-ESP). Funding for GO-ESP was provided by NHLBI grants RC2 HL103010 (HeartGO), RC2 HL102923 (LungGO) and RC2 HL102924 (WHISP). The exome sequencing was performed through NHLBI grants RC2 HL102925 (BroadGO) and RC2 HL102926 (SeattleGO).

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